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AMENDMENTS TO THE CLAIMS

- 1. (Previously Presented) A method of diagnosing a cardiovascular condition characterized by increased expression of a Fit-1/ST2 nucleic acid molecule or an expression product thereof, said method comprising:
- a) contacting a biological sample from a subject with an agent, wherein said agent specifically binds to said Fit-1/ST2 nucleic acid molecule, an expression product thereof, or a fragment of an expression product thereof; and
- b) measuring the amount of bound agent and determining therefrom if the expression of said Fit-1/ST2 nucleic acid molecule or of an expression product thereof is increased relative to a predetermined value, wherein the expression increased relative to a predetermined value is diagnostic of the condition.

2.-5. (Canceled)

- 6. (Previously Presented) The method of claim 1, wherein the cardiovascular condition is selected from the group consisting of myocardial infarction, stroke, arteriosclerosis, and heart failure.
- 7. (Previously Presented) The method of claim 1, wherein the cardiovascular condition is cardiac hypertrophy.
- 8. (Currently amended) A method for monitoring a sample of a patient having or suspected of having cardiovascular condition determining regression, progression or onset of a cardiovascular condition in a subject characterized by increased expression of a nucleic acid molecule or an expression product thereof, comprising:

<u>assaying monitoring</u> a sample from [[a]] <u>the</u> patient[[,]] for increased expression relative to a predetermined value of any of

- (i) a Fit-1/ST2 nucleic acid molecule,
- (ii) a polypeptide encoded by the nucleic acid of part (i), or

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(iii) a peptide fragment of the polypeptide of part (ii).[[,]]

as a determination of the regression, progression or onset of said cardiovascular condition in the subject.

9. (Canceled)

- 10. (Previously Presented) The method of claim 8, wherein the step of monitoring comprises contacting the sample with a detectable agent selected from the group consisting of:
- (a) an isolated nucleic acid molecule which hybridizes to the nucleic acid molecule of part (i), and
- (b) an antibody or an antigen binding fragment thereof which binds the polypeptide of part (ii), or the peptide of part (iii),

11.-36. (Canceled)

- 37. (Previously Presented) The method of claim 1, wherein the sample is a biological fluid or a tissue.
- 38. (Previously Presented) The method of claim 1, wherein the biological fluid is blood or serum.
- 39. (Previously Presented) The method of claim 1, wherein the agent is (i) an isolated nucleic acid molecule that hybridizes to the Fit-1/ST2 nucleic acid molecule or (ii) an antibody that binds the polypeptide encoded by the Fit-1/ST2 nucleic acid molecule, or an antigen-binding fragment of the antibody.
- 40. (Previously Presented) The method of claim 39, wherein the nucleic acid or the antibody is labeled with a radioactive label or an enzyme.

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- 42. (Previously Presented) The method of claim 8, wherein the sample is a biological fluid or a tissue.
- 43. (Previously Presented) The method of claim 8, wherein the biological fluid is blood or serum.
- 44. (Previously Presented) The method of claim 10, wherein the nucleic acid of part (a) or the antibody of part (b) is labeled with a radioactive label or an enzyme.
- 45. (Previously Presented) The method of claim 10, comprising assaying the sample for the peptide of part (iii).
- 46. (Previously Presented) The method of claim 8, wherein the cardiovascular condition is selected from the group consisting of myocardial infarction, stroke, arteriosclerosis, and heart failure.
- 47. (Previously Presented) The method of claim 8, wherein the cardiovascular condition is cardiac hypertrophy.
- 48. (Previously Presented) The method of claim 8, wherein the cardiovascular condition is characterized by mechanical strain, mechanical overload or mechanically-induced deformation in cardiac cells or tissue.